

Y26-D839

Y26-L897

Figure 1



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-255 cgaattcggggcgc  
-240 gtcgaccgcnccagctcgggagacatggggggttaaagctctcgtggnattatcc  
-180 ttcagtggggstattggactgacttttcttatgctgggagtgcccttagaggattatgga  
-120 tttggcagttcacccctgaccatcttgaaaaataagttatctctgatctctgtctgtatgtt  
-60 acttctctccctcaccaacggagaaacaaatgtgggcaaatgtacttctctgaaagtaag  
1 ATGATTTGTCAAAAATTCTGTGTGGTTTGTACATTGGGAATTATTATGTGATAACT  
1 M I C Q K F C V L L H W E F I Y V I T  
61 GCGTTTAACTTGTTCATATCCAATTACTCCTTGGAGATTTAAAGTTGTCTTGCATGCCACCA  
21 A F N L S Y P I T P W R F K L S C M P P  
121 AATTCAACCTATGACTACTTCCTTTTGGCTGCTGGACTCTCAAAGAATACTTCAAATTCTG  
41 N S T Y D Y F L L P A G L S K N T S N S  
A-----A

Figure 2A



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A-----A

181 AATGGACATTATGAGACAGCTGTTGAACCTAAGTTTAATTCAAGTGGTACTCACTTTTCT  
61 N G H Y E T A V E P K F N S S G T H F S

241 AACTTATCCAAACAACTTTCCACTGTTGCTTTTCGGAGTGAGCAAGATAGAAACTGCTCC  
81 N L S K T T F H C C F R S E Q D R N C S

301 TTATGTGCAGACAACATTGAAGGAAGGACATTTGTTTCAACAGTAAATTCTTTAGTTTTT  
101 L C A D N I E G R T F V S T V N S L V F

361 CAACAAATAGATgCAAACTGGAACATACAGTGTGGCTAAAGGAGACTTAAATTTATTC  
121 Q Q I D A N W N I Q C W L K G D L K L F

421 ATCTGTTATGTGGAGTCATTATTAAGAATCTATTCAGGAATTATAACTATAAGGTCAT  
141 I C Y V E S L F K N L F R N Y N Y K V H

B-----B

Figure 2B



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B-----B

481 CTTTATATGTTCTGCCCTGAAGTGTAGAAAGATTACCTCTGGTTCCCCAAAAGGCAGT  
161 L L Y V L P E V L E D S P L V P Q K G S

541 TTTCAGATGGTTCACTGCAATTGCAGTGTTTCATGATGTTGTCAATGTCCTTGTGCCCTGTG  
181 F Q M V H C N C S V H E C C E C L V P V

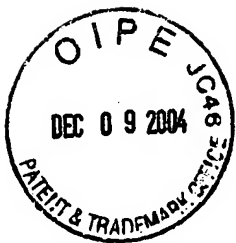
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201 P T A K L N D T L L M C L K I T S G G V

661 ATTTCCrGTCACCTCTAATGTTCAGTTCAGCCCATAAATATGGTGAAGCCTGATCCACCA  
221 I F X S P L M S V Q P I N M V K P D P P

721 TTAGGTTTGCATATGGAAATCACAGATGATGGTAATTTAAAGATTTCTTGGTCCAGCCCA  
241 L G L H M E I T D D G N L K I S W S S P

C-----C

Figure 2C



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C-----C

781 CCATGGTACCATTTCCACTTCAATATCAAGTGAATAATTCAGAGAAATTCACAAACAGTT  
261 P L V P P F P L Q Y Q V K Y S E N S T T V  
841 ATCAGAGAAGCTGACAAAGATTGTCTCAGCTACATCCCTGCTAGTAGACAGTATACTTCCT  
281 I R E A D K I V S A T S L L V D S I L P  
901 GGGTCTTCGTATGAGGTTTCAGGTGAGGGGCAAGAGACTGGATGGCCCAAGGAATCTGGAGT  
301 G S S Y E V Q V R G K R L D G P G I W S  
961 GACTGGAGTACTCCTCGTGTCTTTACCAACACAAGATGTCATATACTTTCCACCTAAAATT  
321 D W S T P R V F T T Q D V I Y F P P K I  
1021 CTGACAAGTGTGGGTCTAATGTTTCTTTTCACTGCACTATAAGAAAGGAAACAAGATT  
341 L T S V G S N V S F H C I Y K K E N K I

D-----D

Figure 2D



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D-----D  
1081 GTTCCCTCAAAAGAGATTGTTGGTGGATGAATTTAGCTGAGAAAATTCCCTCAAAGCCAG  
361 V P S K E I V W M N L A E K I P Q S Q  
1141 TATGATGTTGTGAGTGATCATGTTAGCAAAGTTACTTTTTTCAATCTGAATGAAACCAA  
381 Y D V V S D H V S K V T F F N L N E T K  
1201 CCTCGAGGAAAGTTTACCTATGATGCAGTGTACTGTGCAATGAACATGAATGCCATCAT  
401 P R G K F T Y D A V Y C C N E H E C H H  
1261 CGCTATGCTGAATTATATGTGATTGTGATGTCAATATCAATATCTCATGTGAAACTGATGGG  
421 R Y A E L Y V I D V N I N I S C E T D G  
1321 TACTTAACTAAATGACTTGCAGATGGTCAACCAGTACAATCCAGTCACCTTGCGGAAAGC  
441 Y L T K M T C R W S T S T I Q S L A E S  
E-----E

Figure 2E



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E ----- E  
1381 ACTTTGCAATTGAGGTATCATAGGAGCAGCCCTTTACTGTTCTGATATTCCATCTATTCAT  
461 T L Q L R Y H R S S L Y C S D I P S I H  
1441 CCCATATCTGAGCCCAAAGATTGCTATTTGCAGAGTGATGGTTTTTATGAATGCATTTTC  
481 P I S E P K D C Y L Q S D G F Y A E L C A I F  
1501 CAGCCAAATCTTCCTATTATCTGGCTACACAAATGTGGATTAGGATCAATCACTCTCTAGGT  
501 Q P I F L L S G Y T M W I R I N H S L G  
1561 TCACTTGACTCTCCACCAACAATGTGTCTTCCTGATTCTGTGGTGAAGCCACTGCCCTCCA  
521 S L D S P P T C V L P D S V K P L P P  
1621 TCCAGTGTGAAAGCAGAAATTACTATAAACAATTGGATTATTGAAAAATATCTTGGGAAAAG  
541 S S V K A E I T I N I G L L K I S W E K  
F ----- F

Figure 2F



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F ----- F

1681 CCAGTCTTCCAGAGATAACCTTCAATTCAGATTCCGCTATGGTTTAAGTGGAAGAA  
561 P V F P E N N L Q F Q I R Y G L S G K E

1741 GTACAAATGGAAGATGTATGAGGTTTATGATcCAAaACCAAAATCTGTCAGTCTCCCCAGTT  
581 V Q W K M Y E V Y D P K P K S V S L P V

1801 CCAGACTTGTGTGCAGTCTATGCTGTTCAGGTGCGCTTTAAAGAGGCTAGATGGACTGGGA  
601 P D L C A V Y A V Q V R F K R L D G L G

1861 TATTGGAGTAATTGGAGCAATCCAGCCTACACAGTTGTGATGGATATAAAAGTTCCCTATG  
621 Y W S N W S N P A Y T V M D I K V P M

1921 AGAGGACCTGAATTTGGAGAAATAATTAATGGAGATACTATGAAAAAGGAGAAAAATGTC  
641 R G P E F W R I I N G D T M K K E K N V

G ----- G

Figure 2G





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G ---  
1981 ACTTTACTTTGGAAAGCCCTGATGAAAAATGACTCATGTGCAGTGTTCAGAGATATGTG  
661 T L L W K P L M K N D S L C S V Q R Y V  
2041 ATAAACCATCATACTTCCTSCAATGGAACATGGTCAGAAGATGTGGAAATCACACGAAA  
681 I N H H T S X N G T W S E D V G N H T K  
2101 TTCACTTTCCTGTGGACAGAGCAAGCACATACTGTACGGTCTGGCCATCAATTCAATT  
701 F T F L W T E Q A H T V T V L A I N S I  
2161 GGTGCTTCTGTGCaAATTTTAACCTTTTCATGGCCtATGAGCAAAAGTAAATATC  
721 G A S V A N F N L T F S W P M S K V N I  
2221 GTGCAGTCACTCAGTGCTTATCCTTTAAACAGCAGTTGTGTGATTGTTTCCTGGATACTA  
741 V Q S L S A Y P L N S S C V I V S W I L  
H ---

Figure 2H



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H - - - - - H  
2281 TCACCCAGTGATTACAAGCTAATGTATTTTATTATTGAGTGGAATACTTAATGAAGAT  
761 S P S D Y K L M Y F I I E W K N L N E D  
2341 GGTGAAATAAAATGGCTTAGAATCTCTTCATCTGTTAAGAAAGTATTATATCCATGATCAT  
781 G E I K W L R I S S V K K Y Y I H D H  
2401 TTTATCCCCATTGAGAAGTACCAGTTCAGTCTTTACCCCAATATTATGGGAAGGAGTGGGA  
801 F I P I E K Y Q F S L Y P I F M E G V G  
2461 AAACCAAAGATAATTAAATAGTTTCACTCAAGATGATATTGAAAAACACCAAGATGATGCA  
821 K P K I I N S F T Q D D I E K H Q S D A  
2521 GGTATATGTAAATTGTGCCAGTAATTATTTCCTCTTCCATCTTATTGCTTGGAACATTA  
841 G L Y V I V P V I I S S I L L L G T L  
I - - - - - I

Figure 21

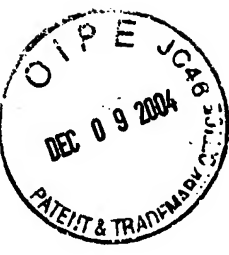


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2581 TTAATATCACACCAAGAATGAAAAAGCTATTTGGGAAGATGTTCCGAACCCCAAGAAT  
861 L I S H Q R M K K L F W E D V P N P K N  
2641 TGTTCTGGGCACAAGGACTTAATTTTCAGAAGAGAACGGACATTCTTTgaaagtctaatac  
881 C S W A Q G L N F Q K R T D I L \*  
2701 atgactactacagatgaacccaatgtgccaaacttcccaacagtctatagagtattagaag  
3761 atttttacattttgaaagaaggaggagcaaatctaaaaaaatttcagttgaaacttctgagag  
2821 ttaacatatgttgattatgttagtaacttaaaatagatgtcatatttaaaccacaagt  
2881 ttacatctaaactcagggtcaaacctcacacactaatttaaaggtttagtagattttcaaat  
2941 ttcatcataagtactaaagaccgaaactaaacagtataaaggaccagatttttgtaattc  
3001 ttttaataccgacaacgacagtaaatgtatagataatttacagtagtttatcatcatctg  
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J.-----J

Figure 2J

|||||



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J.-----J

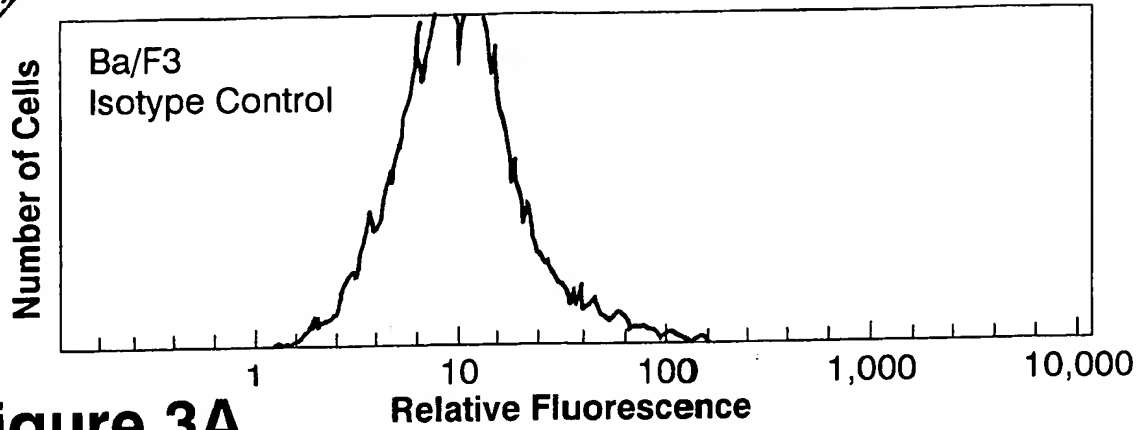
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3301 ataggaaacttttacccaattcactattgaaggcaaaagtcaattttccttcgggcttcaac  
3361 acaaacacgacgggtgtcctgtcacccctcaatgtcaagtatagtcctactgggatgtatg  
3421 ggtccagtctaactgccctgggtcttcccttgtagctgaagattacaggtgcgaaaagaaca  
3481 aattaatactggatttagattaaatgaagggtgacttggtaggttctggagaccgtccgtc  
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Figure 2K

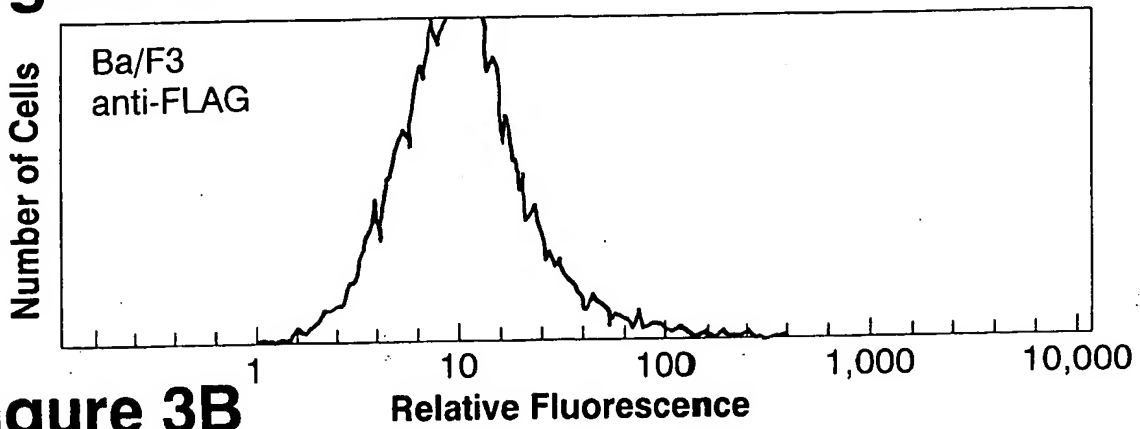




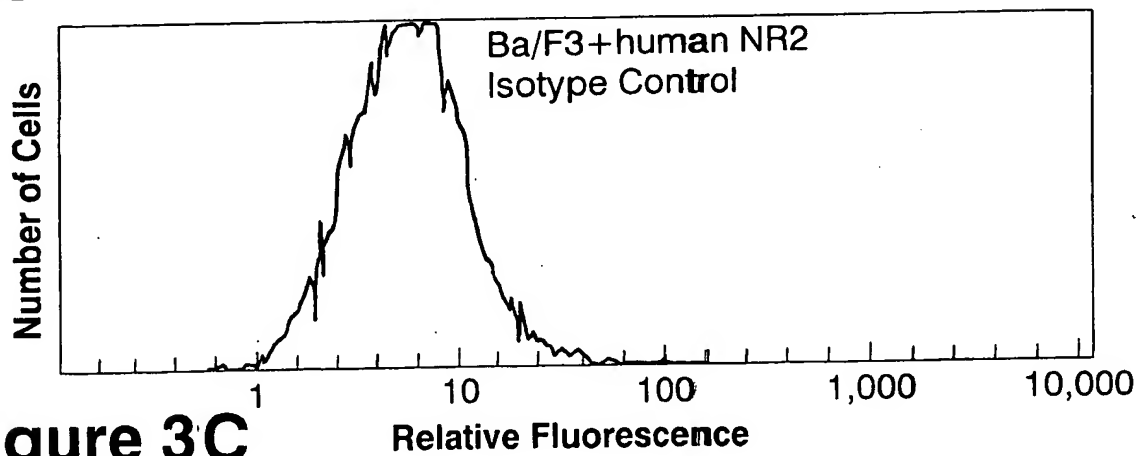
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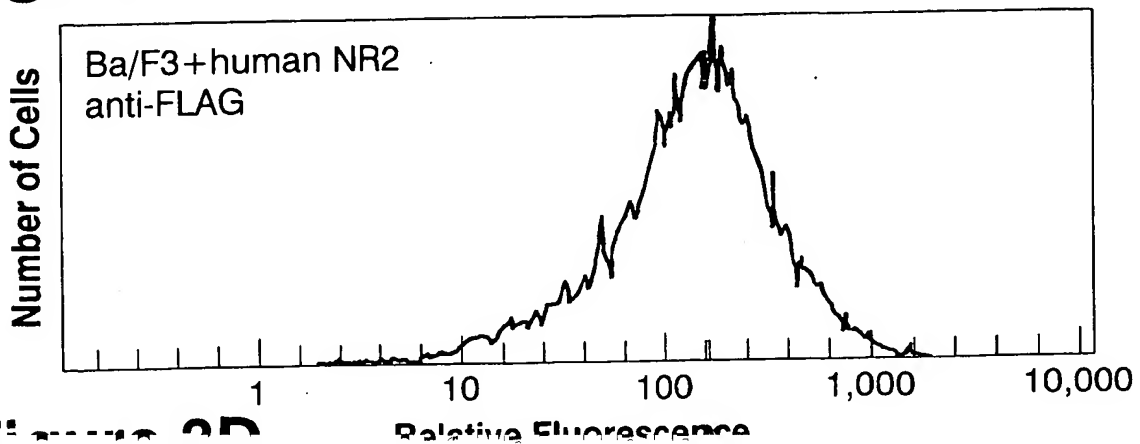
**Figure 3A**



**Figure 3B**

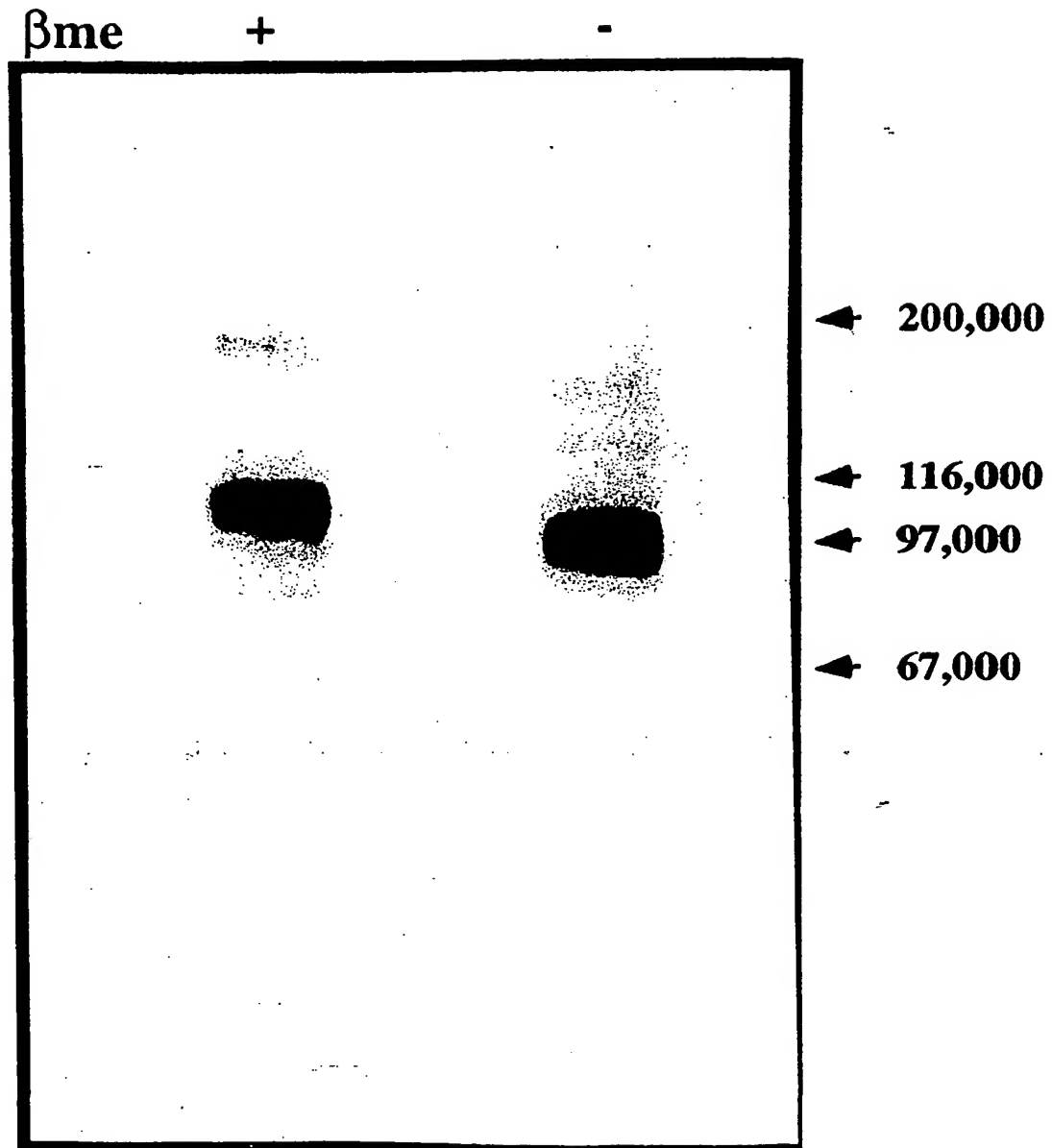


**Figure 3C**





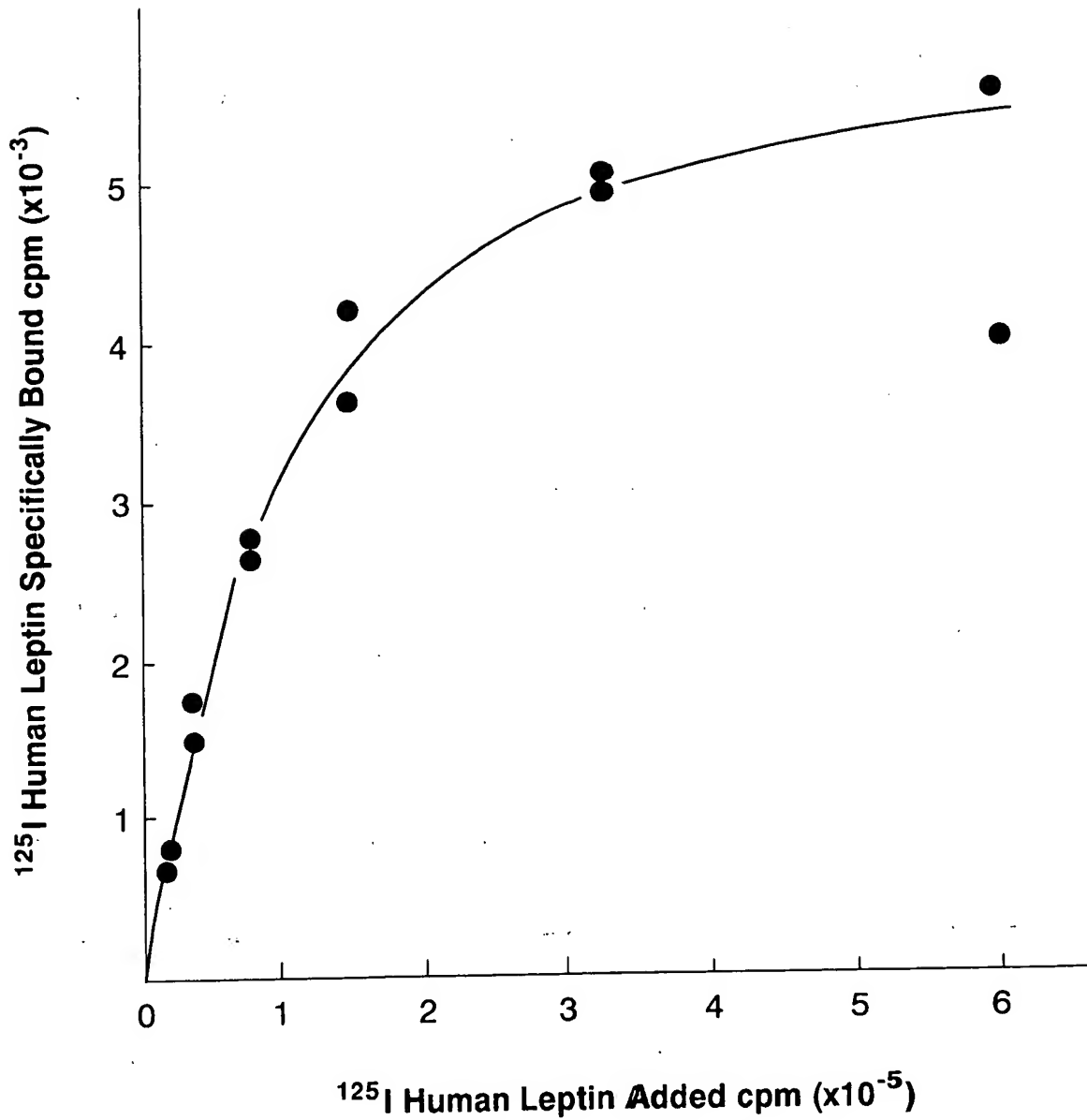
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**Figure 4**



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**Figure 5A**



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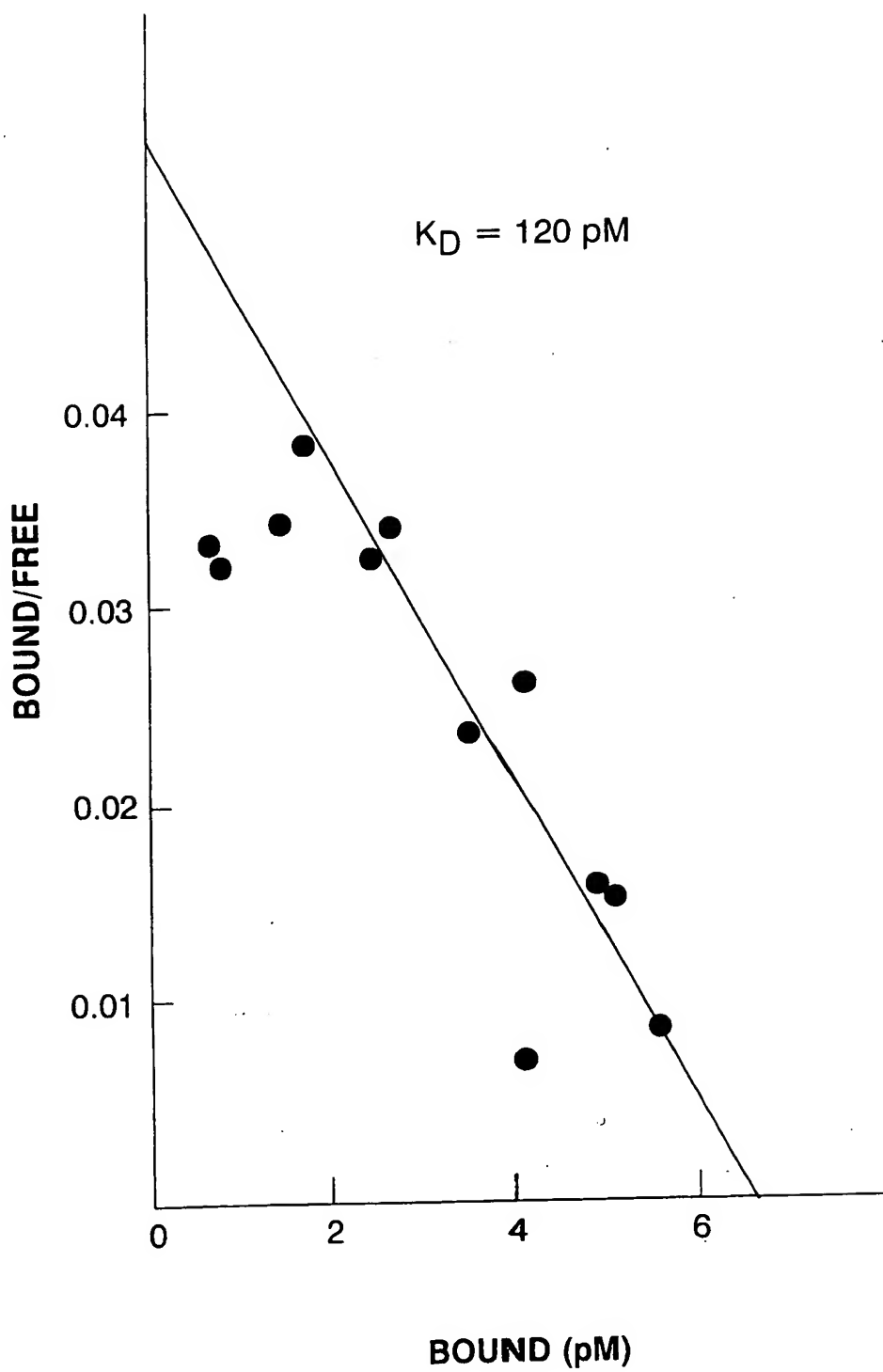


Figure 5B





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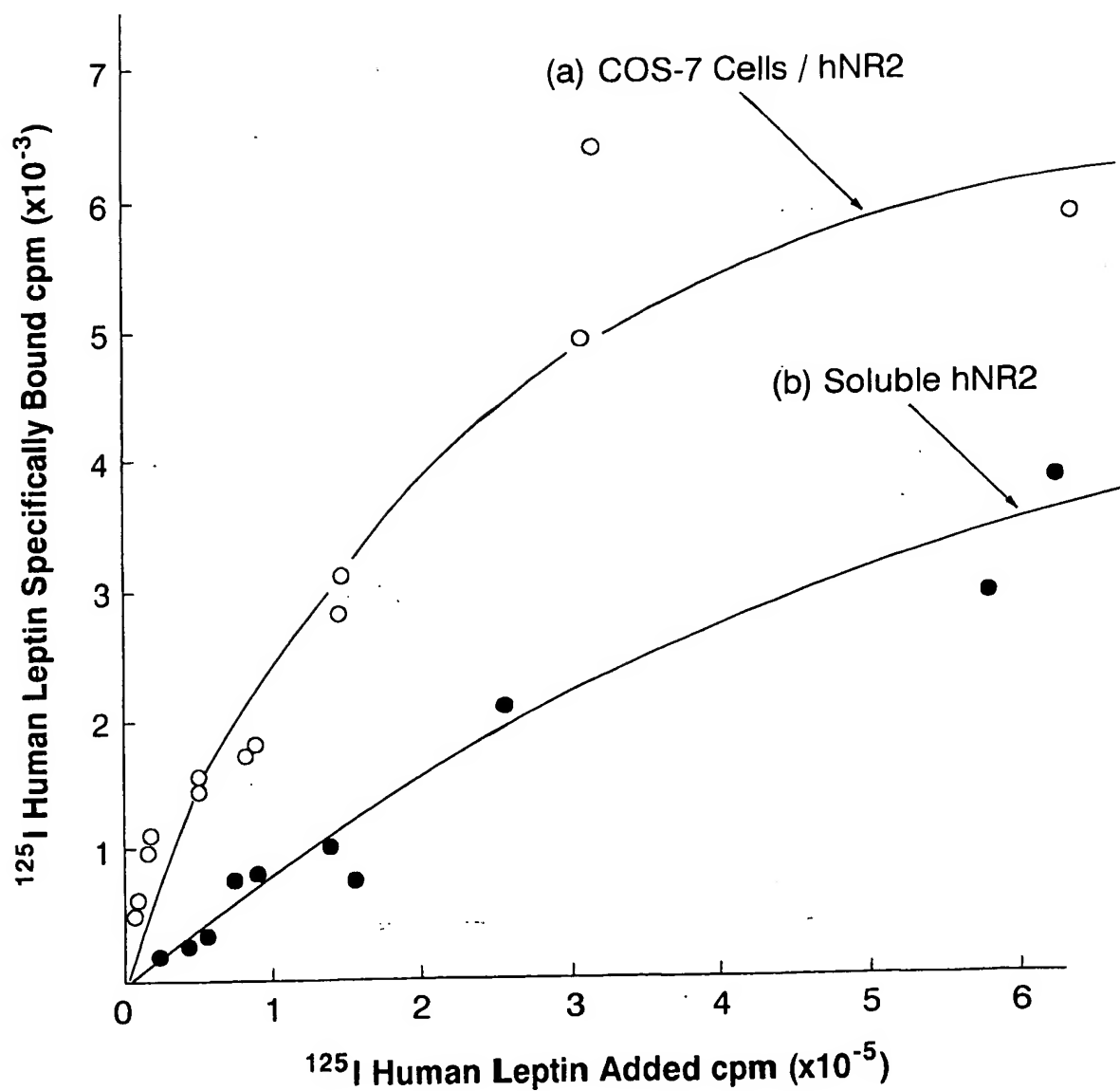
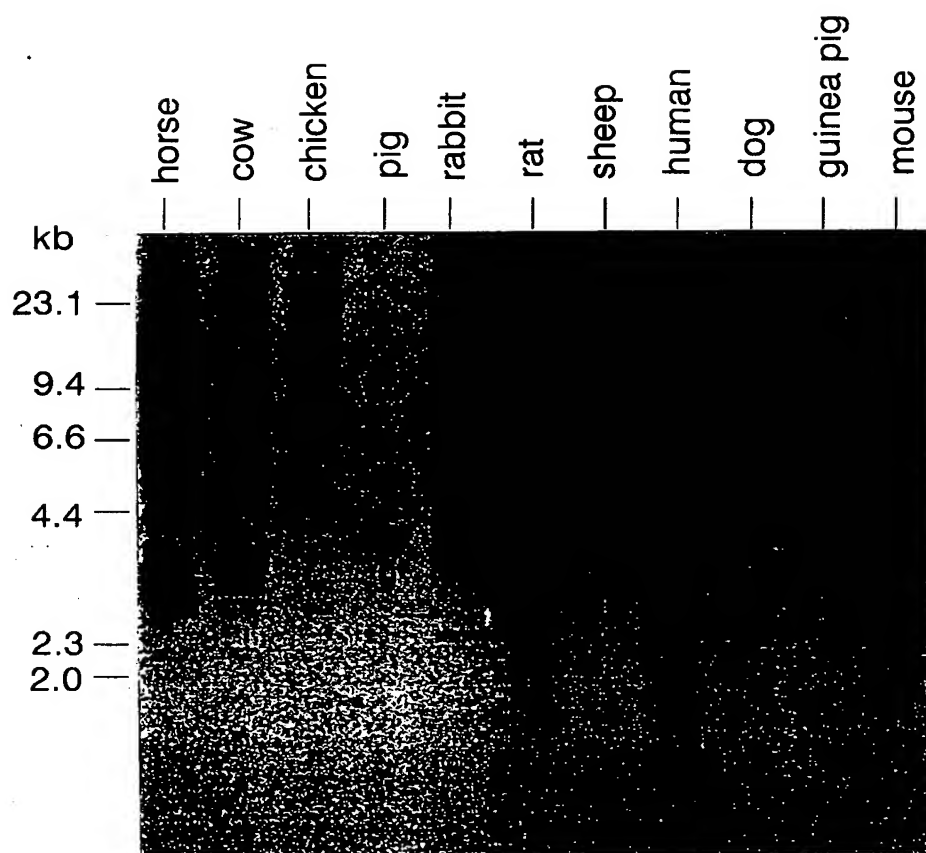


Figure 6



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Cross-species conservation of the NR-2 gene

**Figure 7**



5KB

HAN-1

37-5-6

38-4-1

38-5-2

37-2-1

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EXONS 1 2 3

6

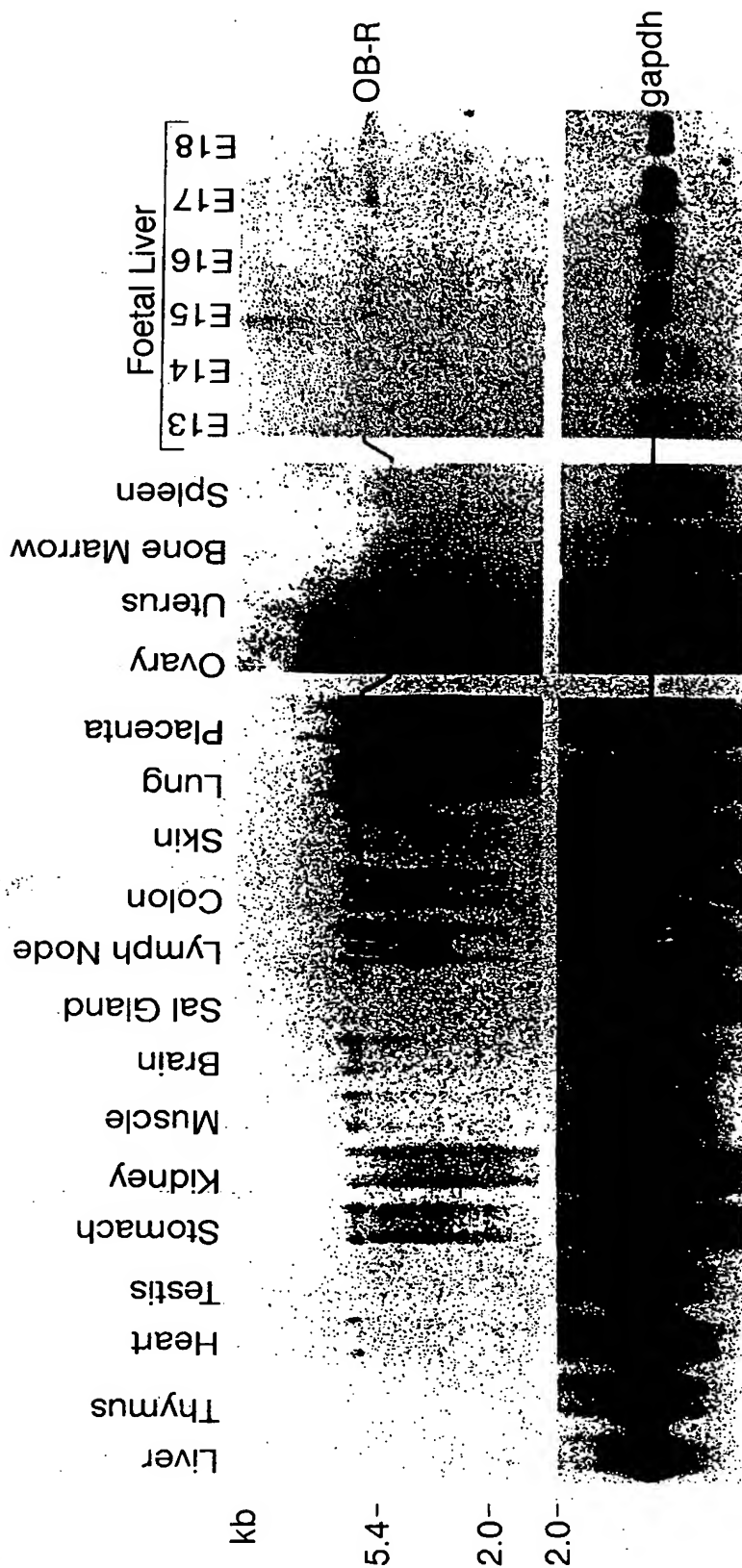
7,8 9

10,11,12

13

15,16,17,18,19

Figure 8



Expression of the Leptin Receptor (NR2) in murine tissues

Figure 9